

IN THE CLAIMS:

Please cancel the current versions of claims 33 and 47 and insert the amended versions of claims 33 and 47 as follows. Pursuant to 37 C.F.R. § 1.121, the following is a clean copy of the amended claims. A marked-up copy of the amended claims is attached hereto on separate sheets.

sub
G2
D1
33. (Twice Amended) A macromolecule comprising a [derivative of a] nucleic acid in isolated form, comprising a fusion of at least two of an oligonucleotide, a polynucleotide and a gene having a nucleotide sequence of at least part of a T-gene selected from the group consisting of the *PLAG* (pleomorphic adenoma gene 1) subfamily of zinc finger protein genes, and at least part of the *CTNNB1* (β catenin) gene and fusions thereof, or complementary or antisense versions of the nucleotide sequence.

D2
sub
G3
47. (Twice Amended) A nucleic acid in isolated form wherein the nucleic acid is one of an oligonucleotide, a polynucleotide and a gene having a sequence of at least part of the *PLAG1* (pleomorphic adenoma gene 1) gene, or the complementary sequence or antisense version of the nucleic acid; wherein a protein encoded by the nucleic acid comprises a polypeptide sequence which is at least 75% identical to a polypeptide sequence of *PLAG 1* in the region from zinc fingers 4 to 7.

Please insert new claims 48 and 49 as follows:

543
H4
D3

48. (New) A macromolecule comprising a nucleic acid in isolated form, comprising a fusion of at least two of an oligonucleotide, a polynucleotide and a gene having a nucleotide sequence of at least part of an intron or exon of the CTNNB1 gene, or the complementary sequence or antisense versions of the nucleotide sequence.

49. (New) A macromolecule according to claim 33 wherein the PLAG subfamily of zinc fingers protein genes comprises the nucleic acids comprising a polypeptide sequence which is at least 75% identical to a polypeptide sequence of PLAG1 in the region from zinc fingers 4 to 7.